SEQUENCE LISTING

<110> Sims, John E.

<120> SIGIRR DNA and Polypeptides

<130> 03260.0044-00304

<140>

<141>

<150> 60/068,770

<151> 1997-12-23

<160> 2

<170> PatentIn Ver. 2.0

<210> 1

<211> 1233

<212> DNA

<213> Homo sapiens

<400> 1

atgccaggtg tctgtgatag ggcccctgac ttcctctccc cgtctgaaga ccaggtgctg 60
aggcctgcct tgggcagctc agtggctctg aactgcacgg cttgggtagt ctctgggccc 120
cactgctccc tgccttcagt ccagtggctg aaagacgggc ttccattggg aattgggggc 180
cactacagcc tccacgagta ctcctgggtc aaggccaacc tgtcagaggt gcttgtgtcc 240
agtgtcctgg gggtcaacgt gaccagcact gaagtctatg gggccttcac ctgctccatc 300
cagaacatca gcttctcctc cttcactctt cagagagctg gccctacaag ccacgtggct 360
gcggtgctgg cctccctcct ggtcctgctg gccctgctgc tggccgccct gctctatgtc 420
aagtgccgtc tcaacgtgct gctctggtac caggacgcgt atggggaggt ggagataaac 480
gacgggaagc tctacgacgc ctacgtctcc tacagcgact gccccgagga ccgcaagttc 540
gtgaacttca tcctaaagcc gcagctggag cggcgtcggg gctacaagct cttcctggac 600
gaccgcgacc tcctgccgcg cgctgagccc tccgccgacc tcttggtgaa cctgagccgc 660
tgccgacgcc tcatcgtggt gctttcggac gccttcctga gccgggcctg gtgcagccac 720
agcttccggg agggcctgtg ccggctgctg gagctcaccc gcagacccat cttcatcacc 780
ttcgagggcc agaggcgcga ccccgcgcac ccggcgctcc gcctgctgcg ccagcaccgc 840

cacctggtga ccttgctgct ctggaggccc ggctccgtga ctccttcctc cgatttttgg 900
aaagaagtgc agctggcgct gccgcggaag gtgcggtaca ggccggtgga aggagacccc 960
cagacgcagc tgcaggacga caaggacccc atgctgattc ttcgaggccg agtccctgag 1020
ggccgggccc tggactcaga ggtggacccg gaccctgagg gcgacctggg tgtccggggg 1080
cctgtttttg gagagccatc agctccaccg cacaccagtg gggtctcgct gggagagagc 1140
cggagcagcg aagtggacgt ctcggatctc ggctcgcgaa actacagtgc ccgcacagac 1200
ttctactgcc tggtgtccaa ggatgatatg tag 1233

<210> 2
<211> 410
<212> PRT
<213> Homo sapiens

<400> 2

Met Pro Gly Val Cys Asp Arg Ala Pro Asp Phe Leu Ser Pro Ser Glu
1 5 10 15

Asp Gln Val Leu Arg Pro Ala Leu Gly Ser Ser Val Ala Leu Asn Cys
20 25 30

Thr Ala Trp Val Val Ser Gly Pro His Cys Ser Leu Pro Ser Val Gln
35 40 45

Trp Leu Lys Asp Gly Leu Pro Leu Gly Ile Gly Gly His Tyr Ser Leu
50 55 60

His Glu Tyr Ser Trp Val Lys Ala Asn Leu Ser Glu Val Leu Val Ser
65 70 75 80

Ser Val Leu Gly Val Asn Val Thr Ser Thr Glu Val Tyr Gly Ala Phe
85 90 95

Thr Cys Ser Ile Gln Asn Ile Ser Phe Ser Ser Phe Thr Leu Gln Arg
100 105 110

Ala Gly Pro Thr Ser His Val Ala Ala Val Leu Ala Ser Leu Leu Val
115 120 125

Leu Leu Ala Leu Leu Leu Ala Ala Leu Leu Tyr Val Lys Cys Arg Leu
130 135 140

Asn Val Leu Leu Trp Tyr Gln Asp Ala Tyr Gly Glu Val Glu Ile Asn
145 150 155 160

Asp Gly Lys Leu Tyr Asp Ala Tyr Val Ser Tyr Ser Asp Cys Pro Glu
165 170 175

Asp Arg Lys Phe Val Asn Phe Ile Leu Lys Pro Gln Leu Glu Arg Arg
180 185 190

Arg Gly Tyr Lys Leu Phe Leu Asp Asp Arg Asp Leu Leu Pro Arg Ala
195 200 205

Glu Pro Ser Ala Asp Leu Leu Val Asn Leu Ser Arg Cys Arg Arg Leu
210 215 220

Ile Val Val Leu Ser Asp Ala Phe Leu Ser Arg Ala Trp Cys Ser His
225 230 235 240

Ser Phe Arg Glu Gly Leu Cys Arg Leu Leu Glu Leu Thr Arg Arg Pro
245 250 255

Ile Phe Ile Thr Phe Glu Gly Gln Arg Arg Asp Pro Ala His Pro Ala
260 265 270

Leu Arg Leu Leu Arg Gln His Arg His Leu Val Thr Leu Leu Leu Trp
275 280 285

Arg Pro Gly Ser Val Thr Pro Ser Ser Asp Phe Trp Lys Glu Val Gln
290 295 300

Leu Ala Leu Pro Arg Lys Val Arg Tyr Arg Pro Val Glu Gly Asp Pro
305 310 315 320

Gln Thr Gln Leu Gln Asp Asp Lys Asp Pro Met Leu Ile Leu Arg Gly
325 330 335

Arg Val Pro Glu Gly Arg Ala Leu Asp Ser Glu Val Asp Pro Asp Pro
340 345 350

Glu Gly Asp Leu Gly Val Arg Gly Pro Val Phe Gly Glu Pro Ser Ala
355 360 365

Pro Pro His Thr Ser Gly Val Ser Leu Gly Glu Ser Arg Ser Ser Glu
370 375 380

Val Asp Val Ser Asp Leu Gly Ser Arg Asn Tyr Ser Ala Arg Thr Asp
385 390 395 400

Phe Tyr Cys Leu Val Ser Lys Asp Asp Met
405 410